

SEP 21 2006

Appl. No.: 10/796,307
Atty. Docket No.: CL1509ORDAMENDMENTS TO THE CLAIMS

This listing of claims will replace all prior versions, and listings of claims in the application.

Listing of claims

1. (Currently amended) A method for identifying an individual who has an altered risk for developing myocardial infarction, comprising detecting a single nucleotide polymorphism (SNP) as represented by a nucleotide sequence selected from the group consisting of SEQ ID NOs 33944, 21614, 36349, 25917, 21749, 29108, 27819, 10810, 11670, 28735 ~~in any one of the nucleotide sequences of SEQ ID NOS:1-450 and 901-43,787~~ in said individual's nucleic acids, wherein the presence of the SNP is correlated with an altered risk for myocardial infarction in said individual.
2. (Original) The method of claim 1 in which the altered risk is an increased risk.
3. (Original) The method of claim 2 in which said individual has previously had a myocardial infarction.
4. (Original) The method of claim 1 in which the altered risk is a decreased risk.
5. (Canceled) The method of claim 1, wherein the SNP is selected from the group consisting of the SNPs set forth in Tables 6-8.
6. (Original) The method of claim 1 in which detection is carried out by a process selected from the group consisting of: allele-specific probe hybridization, allele-specific primer extension, allele-specific amplification, sequencing, 5' nuclease digestion, molecular beacon assay, oligonucleotide ligation assay, size analysis, and single-stranded conformation polymorphism.
7. (Original) An isolated nucleic acid molecule comprising at least 8 contiguous nucleotides wherein one of the nucleotides is a single nucleotide polymorphism (SNP) selected from any one of the nucleotide sequences in SEQ ID NOS:1-450 and 901-43,787, or a complement thereof.

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8. (Original) The isolated nucleic acid molecule of claim 7, wherein the SNP is selected from the group consisting of the SNPs set forth in Tables 3 and 4.

9. (Original) An isolated nucleic acid molecule that encodes any one of the amino acid sequences in SEQ ID NOS:451-900.

10. (Original) An isolated polypeptide comprising an amino acid sequence selected from the group consisting of SEQ ID NOS:451-900.

11. (Original) An antibody that specifically binds to a polypeptide of claim 10, or an antigen-binding fragment thereof.

12. (Original) The antibody of claim 11 in which the antibody is a monoclonal antibody.

13. (Original) An amplified polynucleotide containing a single nucleotide polymorphism (SNP) selected from any one of the nucleotide sequences of SEQ ID NOS:1-450 and 901-43,787, or a complement thereof, wherein the amplified polynucleotide is between about 16 and about 1,000 nucleotides in length.

14. (Original) The amplified polynucleotide of claim 13 in which the nucleotide sequence comprises any one of the nucleotide sequences of SEQ ID NOS:1-450 and 901-43,787.

15. (Original) An isolated polynucleotide which specifically hybridizes to a nucleic acid molecule containing a single nucleotide polymorphism (SNP) in any one of the nucleotide sequences in SEQ ID NOS:1-450 and 901-43,787.

16. (Original) The polynucleotide of claim 15 which is 8-70 nucleotides in length.

17. (Original) The polynucleotide of claim 15 which is an allele-specific probe.

18. (Original) The polynucleotide of claim 15 which is an allele-specific primer.

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19. (Original) The polynucleotide of claim 15, wherein the polynucleotide comprises a nucleotide sequence selected from the group consisting of the primer sequences set forth in Table 5 (SEQ ID NOS:43,788-44,201).

20. (Original) A kit for detecting a single nucleotide polymorphism (SNP) in a nucleic acid, comprising the polynucleotide of claim 15, a buffer, and an enzyme.

21. (Currently amended) A method of detecting a single nucleotide polymorphism (SNP) in a nucleic acid molecule, comprising contacting a test sample with a reagent which specifically hybridizes to a SNP as represented by a nucleotide sequence selected from the group consisting of SEQ ID NOS 33944, 21614, 36349, 25917, 21749, 29108, 27819, 10810, 11670, 28735 ~~in any one of the nucleotide sequences of SEQ ID NOS:1-450 and 901-43,787~~ under stringent hybridization conditions, and detecting the formation of a hybridized duplex.

22. (Original) The method of claim 21 in which detection is carried out by a process selected from the group consisting of: allele-specific probe hybridization, allele-specific primer extension, allele-specific amplification, sequencing, 5' nuclease digestion, molecular beacon assay, oligonucleotide ligation assay, size analysis, and single-stranded conformation polymorphism.

23. (Original) A method of detecting a variant polypeptide, comprising contacting a reagent with a variant polypeptide encoded by a single nucleotide polymorphism (SNP) in any one of the nucleotide sequences of SEQ ID NOS:1-450 and 901-43,787 in a test sample, and detecting the binding of the reagent to the polypeptide.

24. (Original) A method for identifying an agent useful in therapeutically or prophylactically treating myocardial infarction, comprising contacting the polypeptide of claim 10 with a candidate agent under conditions suitable to allow formation of a binding complex between the polypeptide and the candidate agent, and detecting the formation of the binding complex, wherein the presence of the complex identifies said agent.